



1648

RAW SEQUENCE LISTING DATE: 07/12/2002 PATENT APPLICATION: US/09/757,655A TIME: 10:38:42

Input Set : A:\FUJII SEQUENCE LISTING.txt
Output Set: N:\CRF3\07122002\1757655A.raw

ENTERED

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3 <110> APPLICANT: FUJII, Takeru
              YOKOYAMA, Hideakira
              HAMAMOTO, Hidetoshi
      7 <120> TITLE OF INVENTION: A PEPTIDE HAVING AN AFFINITY FOR gp120
      9 <130> FILE REFERENCE: 2001-0019A/LC/01732
     11 <140> CURRENT APPLICATION NUMBER: 09/757,655A
     12 <141> CURRENT FILING DATE: 2001-01-11
     14 <150> PRIOR APPLICATION NUMBER: JP No. 2000-6182
     15 <151> PRIOR FILING DATE: 2000-01-11
     17 <160> NUMBER OF SEQ ID NOS: 19
     19 <170> SOFTWARE: PatentIn Ver. 2.0
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     52 <223> OTHER INFORMATION: Xaa = Gly, Ala, Val, Leu, Ile, Ser, Thr, Met, Asn, Gln, His,
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           Phe, Trp, Pro or Tyr
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    57 `1 . . . 5
    59 <210> SEQ ID NO: 2
    60 <211> LENGTH: 5
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PATENT APPLICATION: US/09/757,655A Input Set : A:\FUJII SEQUENCE LISTING.txt Output Set: N:\CRF3\07122002\I757655A.raw 61 <212> TYPE: PRT 62 <213> ORGANISM: Artificial Sequence 64 <220> FEATURE: 65 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Peptide 67 <220> FEATURE: W--> 68 <221> NAME/KEY: Residue 69 <222> LOCATION: (1) 70 <223> OTHER INFORMATION: Xaa = Asp, Lys, Val, Glu, Gly, Asn or Tyr, or polypeptide residue that an arbitrary amino acid stood in line in the N-terminal side from this amino acid 73 <220> FEATURE: W--> 74 <221> NAME/KEY: Residue 75 <222> LOCATION: (2) 76 <223> OTHER INFORMATION: Xaa = Val, Asp, Trp, Lys, Phe, Ile, Leu or Tyr 78 <220> FEATURE: W--> 79 <221> NAME/KEY: Residue 80 <222> LOCATION: (3) 81 <223> OTHER INFORMATION: Xaa = Lys, Val, Asp, Arg, Ala or Trp 83 <220> FEATURE: W--> 84 <221> NAME/KEY: Residue 85 <222> LOCATION: (4) 86 <223> OTHER INFORMATION: Xaa = Ala, Trp or Gly 88 <220> FEATURE: W--> 89 <221> NAME/KEY: Residue 90 <222> LOCATION: (5) 91 <223> OTHER INFORMATION: Xaa = Gly, Ala, Val, Leu, Ile, Ser, Thr, Met, Asn, Gln, His, 92 Phe, Trp, Pro or Tyr 94 <400> SEQUENCE: 2 / √> 95 Xaa Xaa Xaa Xaa Xaa 98 <210> SEQ ID NO: 3 99 <211> LENGTH: 5 100 <212> TYPE: PRT 101 <213> ORGANISM: Artificial Sequence 103 <220> FEATURE: 104 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Peptide 106 <220> FEATURE: W--> 107 <221> NAME/KEY: Residue 108 <222> LOCATION: (1) 109 <223> OTHER INFORMATION: Xaa = Asp, Lys, Val, Glu, Gly, Asn or Tyr 111 <220> FEATURE: W--> 112 <221> NAME/KEY: Residue 113 <222> LOCATION: (2) 114 <223> OTHER INFORMATION: Xaa = Val, Asp, Trp, Lys, Phe, Ile, Leu or Tyr 116 <220> FEATURE: W--> 117 <221> NAME/KEY: Residue 118 <222> LOCATION: (3) 119 <223> OTHER INFORMATION: Xaa = Lys, Val, Asp, Arg, Ala or Trp

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121 <220> FEATURE: W--> 122 <221> NAME/KEY: Residue

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Input Set : A:\FUJII SEQUENCE LISTING.txt Output Set: N:\CRF3\07122002\I757655A.raw 123 <222> LOCATION: (4) 124 <223> OTHER INFORMATION: Xaa = Ala, Trp or Gly 126 <220> FEATURE: W--> 127 <221> NAME/KEY: Residue 128 <222> LOCATION: (5) 129 <223> OTHER INFORMATION: Xaa = Gly, Ala, Val, Leu, Ile, Thr, Met, Asn, Gln, His, Lys, Arg, Phe, Trp, Pro or Tyr, or polypeptide residue that an arbitrary amino acid 130 stood in line in the C-terminal side of this amino acid, H 133 <400> SEQUENCE: 3 > 134 Xaa Xaa Xaa Xaa Xaa 135 1 138 <210> SEQ ID NO: 4 139 <211> LENGTH: 5 140 <212> TYPE: PRT 141 <213> ORGANISM: Artificial Sequence 143 <220> FEATURE: 144 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Peptide 146 <220> FEATURE: W--> 147 <221> NAME/KEY: Residue 148 <222> LOCATION: (1) 149 <223> OTHER INFORMATION: Xaa = Tyr, Arg, Phe, Gly, Trp, His or Asp 151 <220> FEATURE: W--> 152 <221> NAME/KEY: Residue 153 <222> LOCATION: (2) 154 <223> OTHER INFORMATION: Xaa = Arg, Tyr, Trp, Ala, Val, Gln, His or Lys 156 <220> FEATURE: W--> 157 <221> NAME/KEY: Residue 158 <222> LOCATION: (3) 159 <223> OTHER INFORMATION: Xaa = Lys, Tyr, Arg, Glu, Met or Trp 161 <220> FEATURE: W--> 162 <221> NAME/KEY: Residue 163 <222> LOCATION: (4) 164 <223> OTHER INFORMATION: Xaa = Gly, Ala, Val, Leu, Ile, Ser, Thr, Met, Asn, Gln, His, Lys, Arg, 165 Phe or Trp 167 <220> FEATURE: W--> 168 <221> NAME/KEY: Residue 169 <222> LOCATION: (5) 170 <223> OTHER INFORMATION: Xaa = Gly, Ala, Val, Leu, Ile, Ser, Thr, Met, Asn, Gln, His, Lys, Arg, Phe, Tyr or Trp 171 173 <400× SEQUENCE: 4 174 Xàa Xaa Xaa Xaa Xaa 175 1 177 <210> SEQ ID NO: 5 178 <211> LENGTH: 5 179 <212> TYPE: PRT 180 <213> ORGANISM: Artificial Sequence 182 <220> FEATURE: 183 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Peptide 185 <220> FEATURE:

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Input Set : A:\FUJII SEQUENCE LISTING.txt
Output Set: N:\CRF3\07122002\I757655A.raw

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Input Set : A:\FUJII SEQUENCE LISTING.txt
Output Set: N:\CRF3\07122002\I757655A.raw

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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/757,655A

DATE: 07/12/2002 TIME: 10:38:43

Input Set : A:\FUJII SEQUENCE LISTING.txt
Output Set: N:\CRF3\07122002\I757655A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,5
Seq#:2; Xaa Pos. 1,2,3,4,5
Seq#:3; Xaa Pos. 1,2,3,4,5
Seq#:4; Xaa Pos. 1,2,3,4,5
Seq#:5; Xaa Pos. 1,2,3,4,5
Seq#:6; Xaa Pos. 1,2,3,4,5

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:5; Line(s) 188 Seq#:6; Line(s) 250